

Fig.1

10 20 30 40 50 60  
AACGGATCTG CCCGCCTCAG CCTCCCAAAG TGCTGGGATT GCAGGCGTGA GCCACCTCAC

70 80 90 100 110 120  
CTGGCTACAA GTTTTCAAAA TACATTTATC TAGTACCCAT ACATTCTCCA GTTTGTCCAC

130 140 150 160 170 180  
AGGACATCTT ATGACTTGAG CAAGCTGCTA AAAATCCAAG GGTGCAGCGT TTGTATGTCT

190 200 210 220 230 240  
ATAGGATTGC TCAGATCTGC CCCCACCCTG AAAGAATTTA AGAGAATTTT TTAGGCCAG

250 260 270 280 290 300  
GCACAGTGGC TCACACCTGT AATTCCAGTA CTGTGAGAGT CCGAGGTCAG AGGACTGCTT

PPRE

310 320 330 340 350 360  
GAGGCCAGGA GTTCAAGAGC AGCCTGGACA ACACAGGGAG ACCTGTCACT ACAAAGAATA

370 380 390 400 410 420  
AATAAATTAG CCAGGCTTAG TGGCTCATCC CTGTGGTCCC AGCTACTAGG GAGGCAGAAG

430 440 450 460 470 480  
TAGGACTGCT TGTCCCAGGA GGTCAAGACT GCAGTGAGCT GAGACCCAGC CACCTGCATT

490 500 510 520 530 540  
CCAGCCTGGG CAACAAAAAG AGACCCTGTC TCAAAAAATA AGTTAAATAA ATAAATAATA

550 560 570 580 590 600  
AAAATAGTTT AAACCCTAAA CACATCTTCT TTTTCAAAGA GGA

610 620 630 640 650 660  
GCTGCGTCCT GTTGATCTCC ACTTCCCTTT TTCAGCGTCC ACACTTTTAA CAGTCTCTTT

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Fig.2

670            680            690            700            710            720  
 TGCCAAGGAT AATAAGTATA TAGTTTCTGG AATCCAGATT CTTCCTGTT TGGACAGCCA  
  
 730            740            750            760            770            780  
 GGGGGACAAT TTTTGGTCTG CAGGCCTTTG CATCTGTTCT GCTGTTGCTC AGCAATCTCA  
    GRE  
 790            800            810            820            830            840  
 CAGCAAATTT GCCGAGCCTC TCCGGAATGC ACAGCCAGAC AGAGCTCAGC GCAAAAGCTA  
  
 850            860            870            880            890            900  
 GAGAACCTGG CGGAGGGAGA CTCACAGTGC CACAAAAAAA CTTTATCTTT TCTTTTTTTT  
  
 910            920            930            940            950            960  
 TTTCTTTTCT TTCTTTCTCT TTCTTTCTTG TCTTTCTGTC TTTCCTCTCT CTCTCTCTGT  
  
 970            980            990            1000            1010            1020  
 CTTTCTTTCC TCTCTTTCTT TCTTTTTTCC TACATGGCAA GATCTCCTCA TGGCAGAAAT  
  
 1030            1040            1050            1060            1070            1080  
AATCTGCCTT GACTTCTGTT TCCACGCTGC TTCTGCCAGG ACCATGCGCT CGGCGTGTTT  
    GRE  
 1090            1100            1110            1120            1130            1140  
 TTCTTTCCGC TATAATTATC CAGGCCCATC CCAGCTCTGG TCCCCTCAGC TGTTCCTGG  
  
 1150            1160            1170            1180            1190            1200  
 CAGTCCCTTC TGCTGGTGAA AACACATATG GCGCCGGCCT GACCAGGGTG TAAGTGTGTG  
  
 1210            1220            1230            1240            1250            1260  
 AATATCAGGA AGATGACTGA ACGTCTTTGG GACTCCGTTT CCTCATTGTA AAATGGAGGT

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[illegible]

1270	1280	1290	1300	1310	1320
TAATACCAGC	CTTCTTCTAC	TCCCCAAACG	CACGTGTTTG	TCCCGGCCAG	AGGGCCCAAT
					C/EBP
1330	1340	1350	1360	1370	1380
TGTTGGCTGT	TCACGCATCA	GTTACCCCCA	CAGGACGGGT	CAGCCAATTA	AAGGCCGAACC
				C/EBP	
1390	1400	1410	1420	1430	1440
AGGCCCCGGTC	CATCTCCTGA	CGCCTTTTCT	CATCCCAGGG	CTGGACAGGC	AGCTGGCCTG
				MyoD	
1450	1460	1470	1480	1490	1500
GGCCCCGGCTC	TGCCTTGTCA	CGTGCGGGGG	CCGGCCCCGTT	TGCTTGCTCTG	TGTGTAGGAG
	GRE				
1510	1520	1530	1540	1550	1560
CGTGAGGTCA	CGCTGGGTGC	TCCCGCCCCG	CCGGGGCCTT	TAGTGTCCCT	GGTCCCTAAA
1570	1580	1590	1600	1610	1620
CGCCAGGCCG	CTCCACCGGG	GGAGAAGGCG	CGAACCCAG	CCGAGCCCAA	CGGCTGTTGT
1630	1640	1650	1660	1670	1680
CGGTTGCCGG	GCCACCTGTT	GCTGCAGTTC	TGATTGGTTC	CTTCCCCCGA	CAACGCGGCG
1690	1700	1710	1720	1730	1740
GCTGTAACCA	ATCGACAGCG	AGGCCGGTCG	CGAGGCCCCA	GTCCCGCCCT	GCAGGAGCCA
	C/EBP				
1750	1760	1770	1780	1790	1800
GCCGCGCGCT	CGCTCGCAGG	AGGGTGGGTA	GTTTGCCAG	CGTAGGGGGG	CTGGGCCCCAT
1810	1820	1830	1840	1850	1860
AAAAGAGGAA	GTGCACTTAA	GACACGGCCC	CGCTGGACGC	TTGTTAGAAA	CCGTCCTGGC
1870	1880	1890	1900	1910	1920
TGGGAAGGCA	AGAGGTGTGT	GACTGGACAA	GACTTGTTTC	TGGCGGTCAG	TCTTGCCATC

Fig.4

1930 1940 1950 1960 1970 1980  
CTCACAGAGG TTGGCGGCCC GAGAGAGTGT GAGGCAGAGG CGGGGAGTGG CAAGGGAGTG

1990 2000 2010 2020 2030 2040  
ACCATCTCGG GGAACGAAGG AGTAAACGCG GTGATGGGAC GCACGGAAAC GGGAGTGGAG

2050 2060 2070 2080 2090 2100  
AAAGTCATGG AGAGAACCCT AGGCGGGGCG GTCCCCGCGG AAAGGCGGCT GCTCCAGGGT

2110 2120 2130 2140 2150 2160  
CTCCGCACCC AAGTAGGAGC TGGCAGGCCC GGCCCCGCCC CGCAGGCCCC ACCCCGGGCC

2170 2180 2190 2200 2210 2220  
CCGCCCCCGA GGCTTAAGCC GCGCCGCCGC CTGCGCGGAG CCCCCTGCG AAGCCCAGCT

2230 2240 2250 2260 2270 2280  
GCGCGCGCCT TGGGATTGAC TGTCCACGCT CGCCCGGCTC GTCCGACGCG CCCTCCGCCA

2290 2300 2310 2320 2330 2340  
GCCGACAGAC ACAGCCGCAC GCACTGCCGT GTTCTCCCTG CGGCTCGGTG AGCCTGGCCC

2350 2360 2370 2380 2390 2400  
CAGCCCTGCG CCCTTTGCGC CCCCCACGCT TGTCTGCGT GCGCTGCCCC CTCTTCCATT

2410 2420 2430 2440 2450 2460  
TACCTTCTCT CCCACCCAAG TTTGTACTCT TTTCTTTCTC TCGGTTTTAT TTTTGTGTTT

2470 2480 2490 2500 2510 2520  
TGTTTGTTTG TTTGAGACAG GCTTTCGCTC TGTCTCCAG GCTGGAGTGC AGTGGCGCGA

2530 2540 2550 2560 2570 2580  
TCTCGGCTCA CTGCAGCCTC CACCTCCCAG GTTCAAGCGA TCCGCCTGCC GAGTAGCTGG

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## Fig.5

2590 2600 2610 2620 2630 2640  
GATTACAGGC GCCCGCCACC ACGCCTGGCT AATTTTGTG TTTTGTAGAG ATGGGGTTTC

2650 2660 2670 2680 2690 2700  
GCCATGTTGG CCAGGCTGGC CTCGAAGTGC TCAGCTCAAG CAATCCGCCC GCCTCGGCCT

2710 2720 2730 2740 2750 2760  
CACAAAGTCC TAGAATTTTA GGCATGAGCC TCCGGGTCCG GCCTGTGCTA ATCCTTTCTG

2770 2780 2790 2800 2810 2820  
TCCTTGGTTC TTTATTTCCC TTCTCTCTTT TTCTTAGTCC CTTTGTCTTCT TTCCCTCTCC

2830 2840 2850 2860 2870 2880  
CGTTCAGTTG GCTGTCGTTT GAGCCTCCAC CTTTTCACCTC CCTCCTTTCC ACCACGATGC

2890 2900 2910 2920 2930 2940  
CGAGCCCTGC CTTGGATGGG GACCATCAGC GATGACCACA ATGACCTCTC CCTTACCAGG

2950 2960 2970 2980 2990 3000  
CAGCTCCAGG CAGTGTTCTT GCACCGCCTT TCCCAAGGCT TGGGGGCTTT TTCTAGTGGG

3010 3020 3030 3040 3050 3060  
CTTTGAGCTG CTCAATCTGG CCTCTGCAGG GCCGGCTCCC AGCCCTTCCA ACCTCCTCAC

3070 3080 3090 3100 3110 3120  
AGCCCGACCT GGGACCTAGC CAATTCCCGG AGAGTCTCTG TCCCATCGTG ACCCCCTCAC

3130 3140 3150 3160 3170 3180  
AACTCTCCA CTCACCAAAG TCTGATGACT GTGCTAGGGG GTGCTTATAT AGAGTACTGA

3190 3200 3210 3220 3230 3240  
GTGTTACAAA AGCAGAAGTC TGGATGAGAA CCAATTTGTG ATATTAAGCA GGTGGGGTGG

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Fig.6

3250 3260 3270 3280 3290 3300  
GGGTGGGGAG TGTACCTAGG TTCATTTTCC GCCCTGCTTT TCCCCTTTCC AGTGTGTGCA

3310 3320 3330 3340 3350 3360  
CTTAACCAGT CCCTGGGCCC TGTTCCCAT CCCCCTCCAA GGCATGGATT GGGTGGGCTT

3370 3380 3390 3400 3410 3420  
GTGTGTCTTG GGGCAGGTGG CCCTTTCTAA ACTCTCTGCC TTGCTCACC CACAGGACAC

3430 3440 3450 3460 3470 3480  
ATAGTATGAC CATTAGGTGT TTCGTCTCCC ACCCATTTTC TATGGAAAAC CAAGGGGATC

3490 3500 3510 3520 3530 3540  
GGGCCATGAT AGCCACTGGC AGCTT.....

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Fig.7

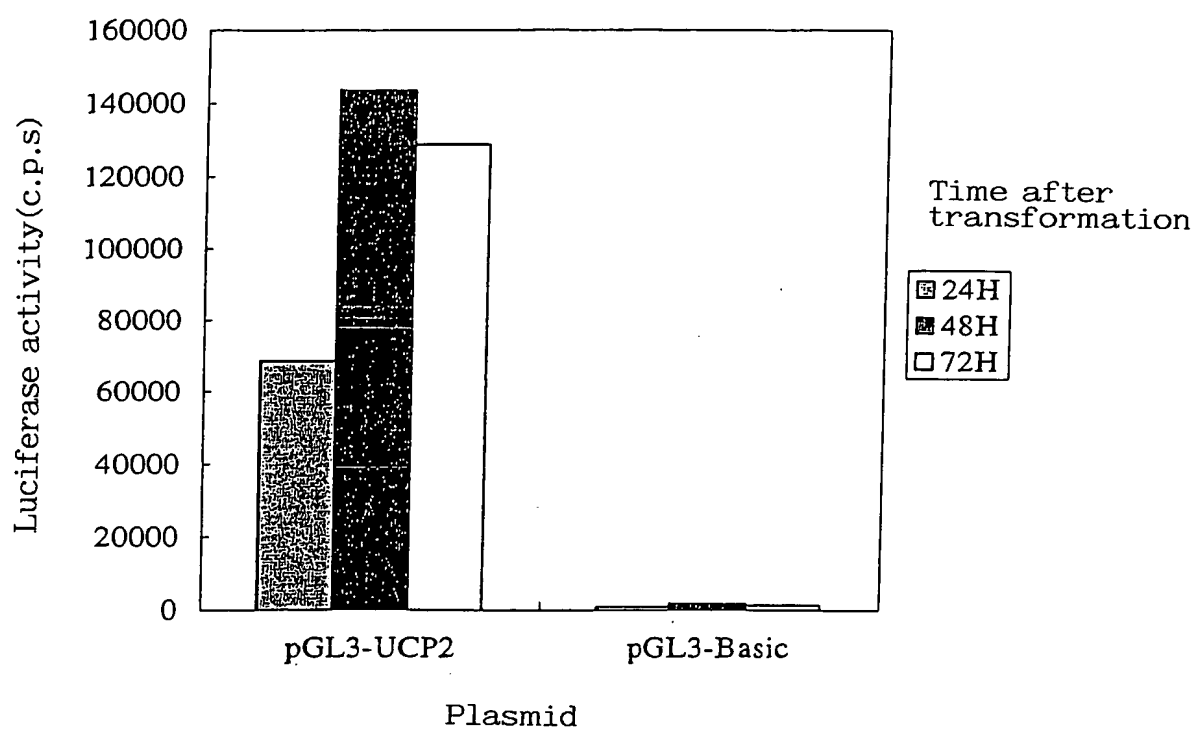


Fig.8

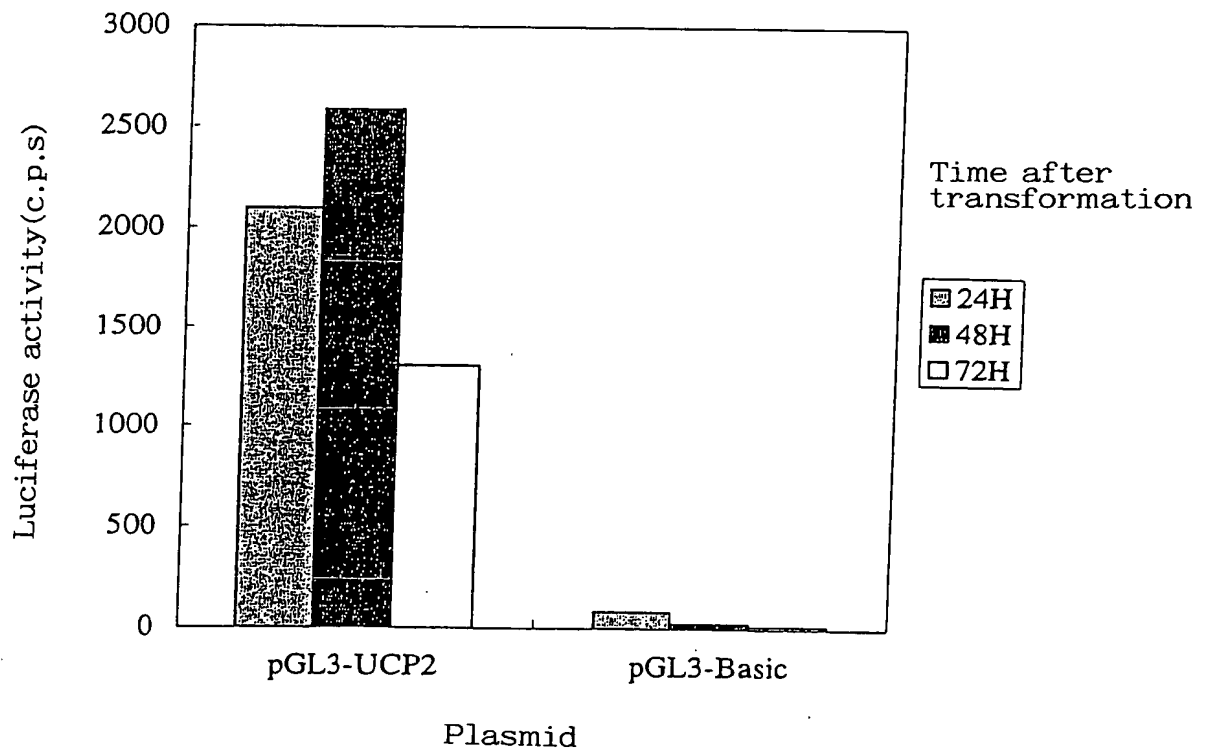




Fig.9

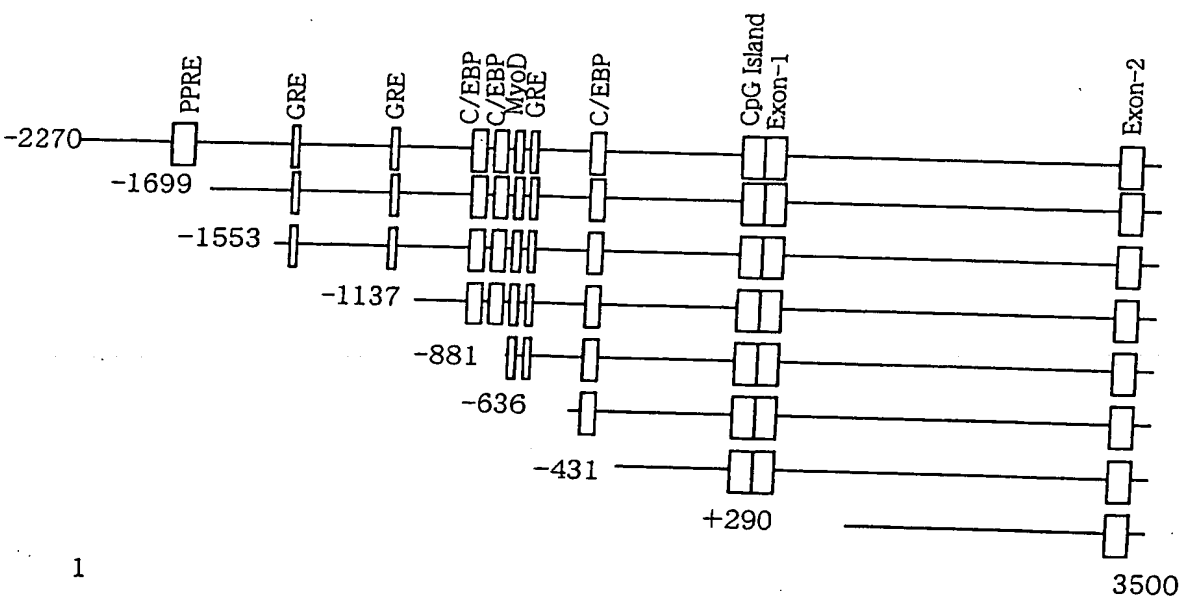
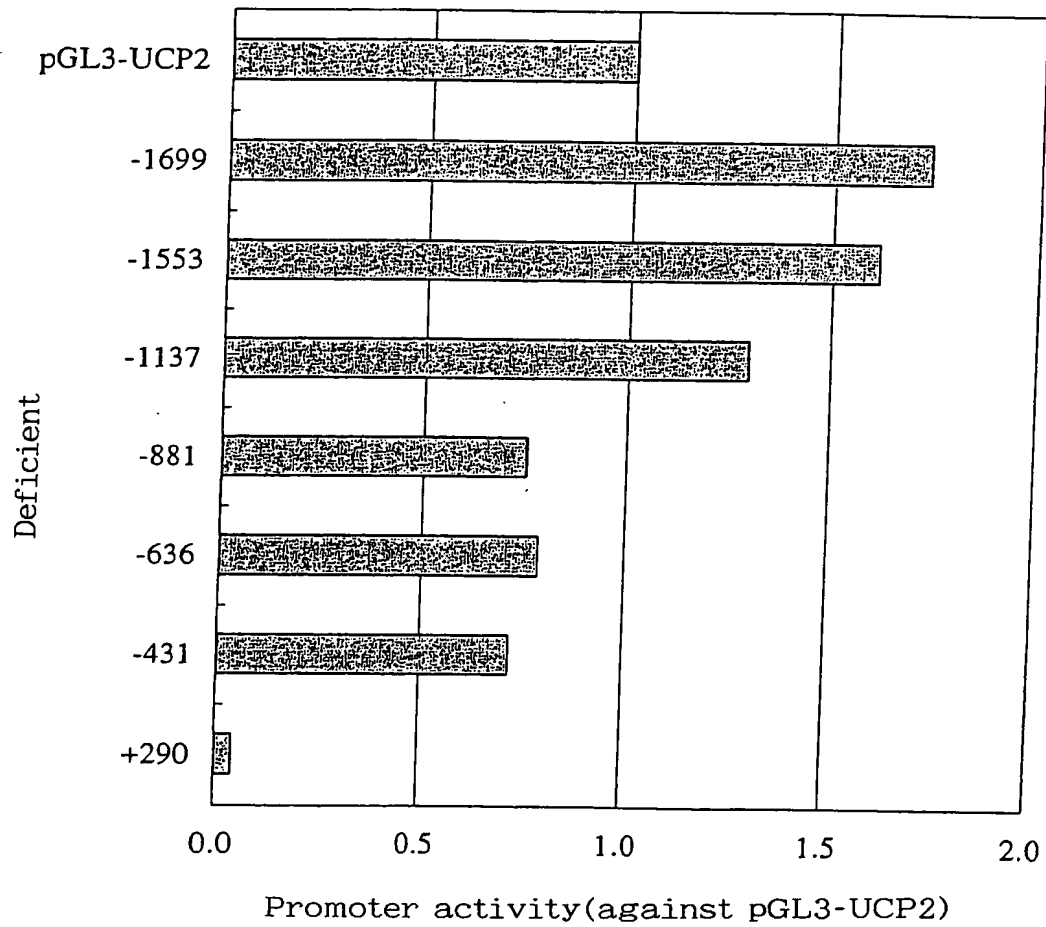


Fig.10



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